

## Amendments to the Specification

Please replace the paragraph at page 11, line 10 with the following replacement paragraph:

The % identity values used herein may be determined by WU-BLAST-2 which was obtained from [Altschul et al., Methods in Enzymology, 266:460-480 (1996); ~~http://blast.wustl.edu/blast/README.html~~ blast.wustl.edu/blast/README.html]. WU-BLAST-2 uses several search parameters, most of which are set to default values. The adjustable parameters are set with the following values: overlap span=1, overlap fraction=0.125, word threshold (T) = 11. The HSPS and HSPS2 parameters are dynamic values and are established by the program itself depending upon the composition of the particular sequence and composition of the particular database against which the sequence of interest is being searched; however, the values may be adjusted to increase sensitivity.